

A standard linear barcode is positioned horizontally across the page, consisting of vertical black bars of varying widths on a white background.

PCT09

ENTERED

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/830,319B

DATE: 03/19/2003
TIME: 14:22:12

Input Set : A:\37921-151292.TXT
Output Set: N:\CRF4\03192003\I830319B.raw

4 <110> APPLICANT: BARTLETT, Perry F.
5 HARTLEY, Lynne
6 PILIZOTTO, Mark
7 KILPATRICK, Trevor
8 KONTGEN, Frank
9 COONAN, Jason
10 GREFERATH, Ursula
11 BOYD, Andrew W.
12 GALEA, Mary
13 PAXINOS, George
14 MURPHY, Mark
16 <120> TITLE OF INVENTION: A Method of Treatment
19 <130> FILE REFERENCE: 37921-151292
21 <140> CURRENT APPLICATION NUMBER: 09/830,319B
22 <141> CURRENT FILING DATE: 2002-10-03
24 <150> PRIOR APPLICATION NUMBER: PCT/AU99/00931
25 <151> PRIOR FILING DATE: 1999-10-27
27 <150> PRIOR APPLICATION NUMBER: PP6748
28 <151> PRIOR FILING DATE: 1998-10-27
30 <160> NUMBER OF SEQ ID NOS: 6
32 <170> SOFTWARE: FastSEQ for Windows Version 4.0
34 <210> SEQ ID NO: 1
35 <211> LENGTH: 27
36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial Sequence
39 <220> FEATURE:
40 <223> OTHER INFORMATION: PCR primer
42 <400> SEQUENCE: 1
43 cgtgctactt ccattttgtca cgtccctg
45 <210> SEQ ID NO: 2
46 <211> LENGTH: 21
47 <212> TYPE: DNA
48 <213> ORGANISM: Artificial Sequence
50 <220> FEATURE:
51 <223> OTHER INFORMATION: PCR primer
53 <400> SEQUENCE: 2
54 tgccgtgata gcaaatttga g
56 <210> SEQ ID NO: 3
57 <211> LENGTH: 22
58 <212> TYPE: DNA
59 <213> ORGANISM: Artificial Sequence
61 <220> FEATURE:
62 <223> OTHER INFORMATION: PCR primer

RAW SEQUENCE LISTING DATE: 03/19/2003
PATENT APPLICATION: US/09/830,319B TIME: 14:22:12

Input Set : A:\37921-151292.TXT
Output Set: N:\CRF4\03192003\I830319B.raw

64 <400> SEQUENCE: 3 22
65 aggaagttag cattatggat ga
67 <210> SEQ ID NO: 4
68 <211> LENGTH: 20
69 <212> TYPE: DNA
70 <213> ORGANISM: Artificial Sequence
72 <220> FEATURE:
73 <223> OTHER INFORMATION: PCR primer
75 <400> SEQUENCE: 4 20
76 tgctcctcggt gcccagcggt
78 <210> SEQ ID NO: 5
79 <211> LENGTH: 27
80 <212> TYPE: DNA
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: PCR primer
86 <400> SEQUENCE: 5 27
87 tttagaattcc ccgaggagga gctgtac
89 <210> SEQ ID NO: 6
90 <211> LENGTH: 27
91 <212> TYPE: DNA
92 <213> ORGANISM: Artificial Sequence
94 <220> FEATURE:
95 <223> OTHER INFORMATION: PCR primer
97 <400> SEQUENCE: 6 27
98 ctagaattct gcagtcccac caccccg

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/830,319B

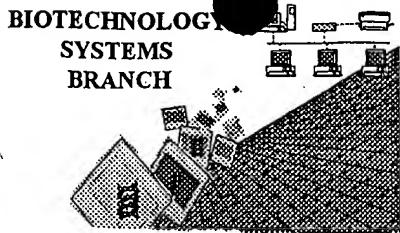
DATE: 03/19/2003

TIME: 14:22:13

Input Set : A:\37921-151292.TXT

Output Set: N:\CRF4\03192003\I830319B.raw

09/8303/9
FY



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/830,319A
Source: PCP/09
Date Processed by STIC: 10/24/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION SERIAL NUMBER: 09/830,319A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleic
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220>
 - - - - - Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



PCT09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/830,319A

DATE: 10/24/2002
TIME: 13:42:19

Input Set : A:\Seqlist.txt
Output Set: N:\CRF4\10242002\I830319A.raw

4 <110> APPLICANT: BARTLETT, Perry F.
5 HARTLEY, Lynne
6 PILIZZOTTO, Mark
7 KILPATRICK, Trevor
8 KONTGEN, Frank
9 COONAN, Jason
10 GREFERATH, Ursula
11 BOYD, Andrew W.
12 GALEA, Mary
13 PAXINOS, George
14 MURPHY, Mark
16 <120> TITLE OF INVENTION: A Method of Treatment
19 <130> FILE REFERENCE: 37921-151292
21 <140> CURRENT APPLICATION NUMBER: 09/830,319A
22 <141> CURRENT FILING DATE: 2002-10-03
24 <150> PRIOR APPLICATION NUMBER: PCT/AU99/00931
25 <151> PRIOR FILING DATE: 1999-10-27
27 <150> PRIOR APPLICATION NUMBER: PP6748
28 <151> PRIOR FILING DATE: 1998-10-27
30 <160> NUMBER OF SEQ ID NOS: 6
32 <170> SOFTWARE: FastSEQ for Windows Version 4.0
34 <210> SEQ ID NO: 1
35 <211> LENGTH: 27
36 <212> TYPE: DNA
37 <213> ORGANISM: Oligonucleotides
39 <400> SEQUENCE: 1
40 cgtgctactt ccattttgtca cgtcctg
42 <210> SEQ ID NO: 2
43 <211> LENGTH: 21
44 <212> TYPE: DNA
45 <213> ORGANISM: Oligonucleotides
47 <400> SEQUENCE: 2
48 tgccgtgata gcaaatttga g
50 <210> SEQ ID NO: 3
51 <211> LENGTH: 22
52 <212> TYPE: DNA
53 <213> ORGANISM: Oligonucleotides
55 <400> SEQUENCE: 3
56 aggaagttag cattatggat ga
58 <210> SEQ ID NO: 4
59 <211> LENGTH: 20
60 <212> TYPE: DNA
61 <213> ORGANISM: Oligonucleotides

Does Not Comply
Corrected Diskette Needed

ppr 1-2

(global error)
invalid response - see item 10 on Error
summary

27 Sheet

21

22

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/830,319A

DATE: 10/24/2002

TIME: 13:42:19

Input Set : A:\Seqlist.txt

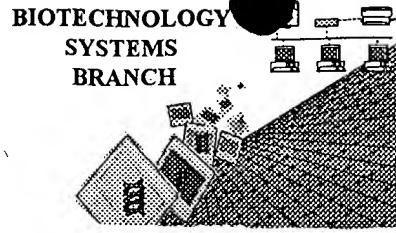
Output Set: N:\CRF4\10242002\I830319A.raw

63 <400> SEQUENCE: 4	
64 tgctcctcgt gcccagcggt	20
66 <210> SEQ ID NO: 5	
67 <211> LENGTH: 27	
68 <212> TYPE: DNA	
69 <213> ORGANISM: Oligonucleotides	
71 <400> SEQUENCE: 5	
72 ttagaattcc ccgaggagga gctgtac	27
74 <210> SEQ ID NO: 6	
75 <211> LENGTH: 27	
76 <212> TYPE: DNA	
77 <213> ORGANISM: Oligonucleotides	
79 <400> SEQUENCE: 6	
80 ctagaattct gcagtccac caccccg	27

VERIFICATION SUMMARY DATE: 10/24/2002
PATENT APPLICATION: US/09/830,319A TIME: 13:42:20

Input Set : A:\Seqlist.txt
Output Set: N:\CRF4\10242002\I830319A.raw

L:22 M:271 C: Current Filing Date differs, Replaced Current Filing Date



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/830,319A
Source: PTO/9
Date Processed by STIC: 10/24/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

Raw Sequence Listing Error Summary

ERROR DETECTED **SUGGESTED CORRECTION** **SERIAL NUMBER: 09/8303/9A**

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleic
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. **Per Sequence Rules**, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220>
- - - - - Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



PCT09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/830,319A

DATE: 10/24/2002
TIME: 13:42:19

Input Set : A:\Seqlist.txt
Output Set: N:\CRF4\10242002\I830319A.raw

4 <110> APPLICANT: BARTLETT, Perry F.
5 HARTLEY, Lynne
6 PILIZZOTTO, Mark
7 KILPATRICK, Trevor
8 KONTGEN, Frank
9 COONAN, Jason
10 GREFERATH, Ursula
11 BOYD, Andrew W.
12 GALEA, Mary
13 PAXINOS, George
14 MURPHY, Mark
16 <120> TITLE OF INVENTION: A Method of Treatment
19 <130> FILE REFERENCE: 37921-151292
21 <140> CURRENT APPLICATION NUMBER: 09/830,319A
22 <141> CURRENT FILING DATE: 2002-10-03
24 <150> PRIOR APPLICATION NUMBER: PCT/AU99/00931
25 <151> PRIOR FILING DATE: 1999-10-27
27 <150> PRIOR APPLICATION NUMBER: PP6748
28 <151> PRIOR FILING DATE: 1998-10-27
30 <160> NUMBER OF SEQ ID NOS: 6
32 <170> SOFTWARE: FastSEQ for Windows Version 4.0
34 <210> SEQ ID NO: 1
35 <211> LENGTH: 27
36 <212> TYPE: DNA
37 <213> ORGANISM: Oligonucleotides (global error)
39 <400> SEQUENCE: 1
40 cgtgctactt ccattttgtca cgtcctg
42 <210> SEQ ID NO: 2
43 <211> LENGTH: 21
44 <212> TYPE: DNA
45 <213> ORGANISM: Oligonucleotides (global error)
47 <400> SEQUENCE: 2
48 tgccgtgata gcaaatttga g
50 <210> SEQ ID NO: 3
51 <211> LENGTH: 22
52 <212> TYPE: DNA
53 <213> ORGANISM: Oligonucleotides (global error)
55 <400> SEQUENCE: 3
56 aggaagttag cattatggat ga
58 <210> SEQ ID NO: 4
59 <211> LENGTH: 20
60 <212> TYPE: DNA
61 <213> ORGANISM: Oligonucleotides (global error)

Does Not Comply
Corrected Diskette Needed

pp 1-2

invalid response - see item 10 on Error
summary

27 Sheet

21

22

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/830,319A

DATE: 10/24/2002

TIME: 13:42:19

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\10242002\I830319A.raw

63 <400> SEQUENCE: 4	
64 tgctcctcgt gcccagcg	20
66 <210> SEQ ID NO: 5	
67 <211> LENGTH: 27	
68 <212> TYPE: DNA	
69 <213> ORGANISM: Oligonucleotides	
71 <400> SEQUENCE: 5	
72 ttagaattcc ccgaggagga gctgtac	27
74 <210> SEQ ID NO: 6	
75 <211> LENGTH: 27	
76 <212> TYPE: DNA	
77 <213> ORGANISM: Oligonucleotides	
79 <400> SEQUENCE: 6	
80 ctagaattct gcagtccac caccccg	27

VERIFICATION SUMMARY DATE: 10/24/2002
PATENT APPLICATION: US/09/830,319A TIME: 13:42:20

Input Set : A:\Seqlist.txt
Output Set: N:\CRF4\10242002\I830319A.raw

L:22 M:271 C: Current Filing Date differs, Replaced Current Filing Date

PATENT
Attorney Docket No.: 37921-151292

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

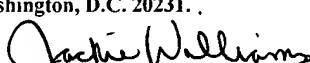
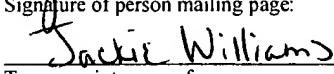
In re: Patent application of :
Perry F. Bartlett, et al. :
Serial No. 09/830,319 : Group Art Unit:
Filing Date: (Int'l.) October 27, 1999 : Examiner:
For: A Method of Treatment :

STATEMENT PURSUANT TO 37 CFR § 1.825

Commissioner for Patents
Box PCT
Washington, D.C. 20231

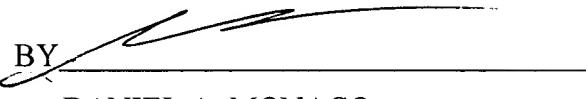
Sir:

The Substitute Sequence Listing filed herewith includes no new matter. The content of the substitute Sequence Listing in computer readable form is the same as the substitute Sequence Listing paper copy submitted herewith.

CERTIFICATE OF MAILING UNDER 37 C.F.R. 1.10	
EXPRESS MAIL Mailing Label Number: <u>EL931089170 US</u>	
Date of Deposit: <u>11/21/02</u>	
I hereby certify that this correspondence, along with any paper referred to as being attached or enclosed, and/or fee, is being deposited with the United States Postal Service, "EXPRESS MAIL-POST OFFICE TO ADDRESSEE" service under 37 CFR 1.10, on the date indicated above, and addressed to: Commissioner for Patents, Washington, D.C. 20231.	
 Signature of person mailing page:	
 Type or print name of person	

Respectfully submitted,

PERRY F. BARTLETT, et al

BY 

DANIEL A. MONACO
Reg. No. 30,480
DRINKER, BIDDLE & REATH, LLP.
One Logan Square
18th and Cherry Streets
Philadelphia, PA 19103-6996
(215) 988-3312
(215) 988-2757 – fax
Attorney for the Applicants